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H-7

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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/040,394

DATE: 07/09/2002  
TIME: 16:10:45

Input Set : A:\PTO.VSK.TXT  
Output Set: N:\CRF3\07092002\J040394.raw

ENTERED

3 <110> APPLICANT: Soe, Jorn  
4 Poulsen, Charlotte  
5 Rasmussen, Preben  
6 Madrid, Susan  
7 Zargahi, Masoud  
9 <120> TITLE OF INVENTION: Improved Method for Preparing Flour Doughs and Products Made  
From Such

10 Doughs Using a Glycerol Oxidase  
12 <130> FILE REFERENCE: 674509-2045.1  
14 <140> CURRENT APPLICATION NUMBER: US 10/040,394  
C--> 15 <141> CURRENT FILING DATE: 2002-06-21  
17 <150> PRIOR APPLICATION NUMBER: US 09/402,664  
18 <151> PRIOR FILING DATE: 1998-04-03  
20 <150> PRIOR APPLICATION NUMBER: PCT/DK98/00136  
21 <151> PRIOR FILING DATE: 1998-04-03  
23 <150> PRIOR APPLICATION NUMBER: DK 0400/97  
24 <151> PRIOR FILING DATE: 1997-04-09  
26 <160> NUMBER OF SEQ ID NOS: 22  
28 <170> SOFTWARE: PatentIn version 3.1  
30 <210> SEQ ID NO: 1  
31 <211> LENGTH: 25  
32 <212> TYPE: PRT  
33 <213> ORGANISM: Aspergillus tubingensis  
35 <220> FEATURE:  
36 <221> NAME/KEY: MISC\_FEATURE  
37 <222> LOCATION: (22)..(22)  
38 <223> OTHER INFORMATION: "Xaa" can be any amino acid  
41 <400> SEQUENCE: 1  
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44 1 5 10 15

W--> 47 Ser Ala Ala Ala Tyr Xaa Ser Asn Asn

48 20 25  
51 <210> SEQ ID NO: 2  
52 <211> LENGTH: 7  
53 <212> TYPE: PRT  
54 <213> ORGANISM: Aspergillus tubingensis  
56 <400> SEQUENCE: 2  
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59 1 5  
62 <210> SEQ ID NO: 3  
63 <211> LENGTH: 14  
64 <212> TYPE: PRT  
65 <213> ORGANISM: Aspergillus tubingensis  
67 <400> SEQUENCE: 3

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69 Ala Trp Glu Ser Ala Ala Asp Glu Leu Thr Ser Lys Ile Lys  
70 1 5 10  
73 <210> SEQ ID NO: 4  
74 <211> LENGTH: 20  
75 <212> TYPE: DNA  
76 <213> ORGANISM: Artificial Sequence  
78 <220> FEATURE:  
79 <223> OTHER INFORMATION: PCR primer used for PCR amplification of a fragment of the  
lipase  
80 gene  
82 <220> FEATURE:  
83 <221> NAME/KEY: misc\_feature  
84 <222> LOCATION: (9)..(9)  
85 <223> OTHER INFORMATION: "n" can be a or t/u or g or c  
88 <220> FEATURE:  
89 <221> NAME/KEY: misc\_feature  
90 <222> LOCATION: (12)..(12)  
91 <223> OTHER INFORMATION: "n" can be a or t/u or g or c  
94 <220> FEATURE:  
95 <221> NAME/KEY: misc\_feature  
96 <222> LOCATION: (18)..(18)  
97 <223> OTHER INFORMATION: "n" can be a or t/u or g or c  
100 <400> SEQUENCE: 4  
  
Wt-> 101 ttccaraanc cngtrtgnac 20  
104 <210> SEQ ID NO: 5  
105 <211> LENGTH: 18  
106 <212> TYPE: DNA  
107 <213> ORGANISM: Artificial Sequence  
109 <220> FEATURE:  
110 <223> OTHER INFORMATION: PCR primer used for PCR amplification of a fragment of the  
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111 gene  
113 <220> FEATURE:  
114 <221> NAME/KEY: misc\_feature  
115 <222> LOCATION: (6)..(6)  
116 <223> OTHER INFORMATION: "n" can be a or t/u or g or c  
119 <220> FEATURE:  
120 <221> NAME/KEY: misc\_feature  
121 <222> LOCATION: (12)..(12)  
122 <223> OTHER INFORMATION: "n" can be a or t/u or g or c  
125 <400> SEQUENCE: 5  
  
Wt-> 126 carytnnttyg cncartgg 18  
129 <210> SEQ ID NO: 6  
130 <211> LENGTH: 17  
131 <212> TYPE: DNA  
132 <213> ORGANISM: Artificial Sequence  
134 <220> FEATURE:  
135 <223> OTHER INFORMATION: PCR primer used for PCR amplification of a fragment of the  
lipase  
136 gene  
138 <400> SEQUENCE: 6  
139 gcvgchswyt cccavgc 17

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Input Set : A:\PTO.VSK.TXT  
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142 <210> SEQ ID NO: 7  
 143 <211> LENGTH: 317  
 144 <212> TYPE: DNA  
 145 <213> ORGANISM: Aspergillus tubingensis  
 147 <400> SEQUENCE: 7

148 cagttgttcg cgcaatggtc tgccgcagct tattgctcga ataatatcga ctgcggaaagav 60  
 150 tccaaacttga catgcacggc caacgcctgt ccatcagtcg aggaggccag taccacgatg 120  
 152 ctgctggagt tcgaccctgta tgtcaactcag atcgcagaca tagagcacag ctaattgaac 180  
 154 aqgacgaacg acttttggag gcacagccgg tttcctggcc gcggacaaca ccaacaagcgc 240  
 156 gctcggttc gccttccggg gaagcagcac gattgagaac tggattgcta atcytgactt 300  
 158 catcctggra gataacg 317

161 <210> SEQ ID NO: 8  
 162 <211> LENGTH: 1045  
 163 <212> TYPE: DNA  
 164 <213> ORGANISM: Aspergillus tubingensis  
 166 <400> SEQUENCE: 8

167 atgttctctg gacgggttgg agtgctttg acagcgcttg ctgcgtggg tgctgccgcg 60  
 169 ccggcaccgc ttgctgtcg gagtaggtgt gcccgtatgt agatgggtgg atagoactga 120  
 171 tqaagggtga atagggtgtct cgacttccac gttggatgag ttgcaattgt tcgcgcaatg 180  
 173 gtctgccca gcttattgtct cgaataatat cgactcgaaa gactccaact tgacatgcac 240  
 175 ggcacaacgccc tgtccatcag tcgaggaggc cagtaccacg atgctgtcg agttcgcac 300  
 177 gtatgtcaact cagatcgacg acatagagca cagctaattt gaacaggacg aacgactttg 360  
 179 gaggcacago cggtttctg gccgcggaca acaccaacaa gcccgtcg tgccgttcc 420  
 181 ggggaagcagc cacgattgag aactggattt ctaatcttgc cttcatcctg gaagataacg 480  
 183 acgacctctg caccggctgc aaggccata ctgggttctg gaaggcatgg gactccgtg 540  
 185 ccgacgaact gacgagaacg atcaagtctg cgatgagcac gtattcgggc tataccctat 600  
 187 acttcaccgg gcacagttt ggcggcgcatttggctacgc gggagcgaca gttctgcgaa 660  
 189 atgacggata tagcgttgag ctggtgagtc cttcacaaag gtgtatggagc gacaatcggg 720  
 191 aacagacagt caatagatac cctatggatg tcctcaatc gggaaactatg cgctggctga 780  
 193 gcatatcacc agtcaggat ctggggccaa cttccgtt acacacttgc acgacatcg 840  
 195 cccccgggtg ccacccatgg actttggatt cagtcagccaa agtccggaaat actggatcac 900  
 197 cagtggcaat ggagccagtg tcacggcgtc ggatatcgaa gtcatcgagg gaatcaattc 960  
 199 aacggcgggaa aatgcaggcg aagcaacggc gagcgttgc gtcacttgc ggtactttt 1020  
 201 tqcgatttcc gagtgcgtc tataa 1045

204 <210> SEQ ID NO: 9  
 205 <211> LENGTH: 297  
 206 <212> TYPE: PRT  
 207 <213> ORGANISM: Aspergillus tubingensis  
 209 <400> SEQUENCE: 9

211 Met Phe Ser Gly Arg Phe Gly Val Leu Leu Thr Ala Leu Ala Ala Leu  
 212 1 5 10 15  
 215 Gly Ala Ala Ala Pro Ala Pro Leu Ala Val Arg Ser Val Ser Thr Ser  
 216 20 25 30  
 219 Thr Leu Asp Glu Leu Gln Leu Phe Ala Gln Trp Ser Ala Ala Ala Tyr  
 220 35 40 45  
 223 Cys Ser Asn Asn Ile Asp Ser Lys Asp Ser Asn Leu Thr Cys Thr Ala  
 224 50 55 60  
 227 Asn Ala Cys Pro Ser Val Glu Glu Ala Ser Thr Thr Met Leu Leu Glu  
 228 65 70 75 80

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231 Phe Asp Leu Thr Asn Asp Phe Gly Gly Thr Ala Gly Phe Leu Ala Ala  
232 85 90 95  
235 Asp Asn Thr Asn Lys Arg Leu Val Val Ala Phe Arg Gly Ser Ser Thr  
236 100 105 110  
239 Ile Glu Asn Trp Ile Ala Asn Leu Asp Phe Ile Leu Glu Asp Asn Asp  
240 115 120 125  
243 Asp Leu Cys Thr Gly Cys Lys Val His Thr Gly Phe Trp Lys Ala Trp  
244 130 135 140  
247 Glu Ser Ala Ala Asp Glu Leu Thr Ser Lys Ile Lys Ser Ala Met Ser  
248 145 150 155 160  
251 Thr Tyr Ser Gly Tyr Thr Leu Tyr Phe Thr Gly His Ser Leu Gly Gly  
252 165 170 175  
255 Ala Leu Ala Thr Leu Gly Ala Thr Val Leu Arg Asn Asp Gly Tyr Ser  
256 180 185 190  
259 Val Glu Leu Tyr Thr Tyr Gly Cys Pro Arg Ile Gly Asn Tyr Ala Leu  
260 195 200 205  
263 Ala Glu His Ile Thr Ser Gln Gly Ser Gly Ala Asn Phe Arg Val Thr  
264 210 215 220  
267 His Leu Asn Asp Ile Val Pro Arg Val Pro Pro Met Asp Phe Gly Phe  
268 225 230 235 240  
271 Ser Gln Pro Ser Pro Glu Tyr Trp Ile Thr Ser Gly Asn Gly Ala Ser  
272 245 250 255  
275 Val Thr Ala Ser Asp Ile Glu Val Ile Glu Gly Ile Asn Ser Thr Ala  
276 260 265 270  
279 Gly Asn Ala Gly Glu Ala Thr Val Ser Val Val Ala His Leu Trp Tyr  
280 275 280 285  
283 Phe Phe Ala Ile Ser Glu Cys Leu Leu  
284 290 295  
287 <210> SEQ ID NO: 10  
288 <211> LENGTH: 392  
289 <212> TYPE: PRT  
290 <213> ORGANISM: Rhizopus delamar  
292 <400> SEQUENCE: 10  
294 Met Val Ser Phe Ile Ser Ile Ser Gln Gly Val Ser Leu Cys Leu Leu  
295 1 5 10 15  
298 Val Ser Ser Met Met Leu Gly Ser Ser Ala Val Pro Val Ser Gly Lys  
299 20 25 30  
302 Ser Gly Ser Ser Asn Thr Ala Val Ser Ala Ser Asp Asn Ala Ala Leu  
303 35 40 45  
306 Pro Pro Leu Ile Ser Ser Arg Cys Ala Pro Pro Ser Asn Lys Gly Ser  
307 50 55 60  
310 Lys Ser Asp Leu Gln Ala Glu Pro Tyr Asn Met Gln Lys Asn Thr Glu  
311 65 70 75 80  
314 Trp Tyr Glu Ser His Gly Gly Asn Leu Thr Ser Ile Gly Lys Arg Asp  
315 85 90 95  
318 Asp Asn Leu Val Gly Gly Met Thr Leu Asp Leu Pro Ser Asp Ala Pro  
319 100 105 110  
322 Pro Ile Ser Leu Ser Ser Ser Thr Asn Ser Ala Ser Asp Gly Gly Lys  
323 115 120 125

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326 Val Val Ala Ala Thr Thr Ala Gln Ile Gln Glu Phe Thr Lys Tyr Ala  
 327 130 135 140  
 330 Gly Ile Ala Ala Thr Ala Tyr Cys Arg Ser Val Val Pro Gly Asn Lys  
 331 145 150 155 160  
 334 Trp Asp Cys Val Gln Cys Gln Lys Trp Val Pro Asp Gly Lys Ile Ile  
 335 165 170 175  
 338 Thr Thr Phe Thr Ser Leu Leu Ser Asp Thr Asn Gly Tyr Val Leu Arg  
 339 180 185 190  
 342 Ser Asp Lys Gln Lys Thr Ile Tyr Leu Val Phe Arg Gly Thr Asn Ser  
 343 195 200 205  
 346 Phe Arg Ser Ala Ile Thr Asp Ile Val Phe Asn Phe Ser Asp Tyr Lys  
 347 210 215 220  
 350 Pro Val Lys Gly Ala Lys Val His Ala Gly Phe Leu Ser Ser Tyr Glu  
 351 225 230 235 240  
 354 Gln Val Val Asn Asp Tyr Phe Pro Val Val Gln Glu Gln Leu Thr Ala  
 355 245 250 255  
 358 His Pro Thr Tyr Lys Val Ile Val Thr Gly His Ser Leu Gly Gly Ala  
 359 260 265 270  
 362 Gln Ala Leu Leu Ala Gly Met Asp Leu Tyr Gln Arg Glu Pro Arg Leu  
 363 275 280 285  
 366 Ser Pro Lys Asn Leu Ser Ile Phe Thr Val Gly Gly Pro Arg Val Gly  
 367 290 295 300  
 370 Asn Pro Thr Phe Ala Tyr Tyr Val Glu Ser Thr Gly Ile Pro Phe Gln  
 371 305 310 315 320  
 374 Arg Thr Val His Lys Arg Asp Ile Val Pro His Val Pro Pro Gln Ser  
 375 325 330 335  
 378 Phe Gly Phe Leu His Pro Gly Val Glu Ser Trp Ile Lys Ser Gly Thr  
 379 340 345 350  
 382 Ser Asn Val Gln Ile Cys Thr Ser Glu Ile Glu Thr Lys Asp Cys Ser  
 383 355 360 365  
 386 Asn Ser Ile Val Pro Phe Thr Ser Ile Leu Asp His Leu Ser Tyr Phe  
 387 370 375 380  
 390 Asp Ile Asn Glu Gly Ser Cys Leu  
 391 385 390  
 394 <210> SEQ ID NO: 11  
 395 <211> LENGTH: 363  
 396 <212> TYPE: PRT  
 397 <213> ORGANISM: Rhizomucor miehei  
 399 <400> SEQUENCE: 11

401 Met Val Leu Lys Gln Arg Ala Asn Tyr Leu Gly Phe Leu Ile Val Phe  
 402 1 5 10 15  
 405 Phe Thr Ala Phe Leu Val Glu Ala Val Pro Ile Lys Arg Gln Ser Asn  
 406 20 25 30  
 409 Ser Thr Val Asp Ser Leu Pro Pro Leu Ile Pro Ser Arg Thr Ser Ala  
 410 35 40 45  
 413 Pro Ser Ser Ser Pro Ser Thr Thr Asp Pro Glu Ala Pro Ala Met Ser  
 414 50 55 60  
 417 Arg Asn Gly Pro Leu Pro Ser Asp Val Glu Thr Lys Tyr Gly Met Ala  
 418 65 70 75 80

RAW SEQUENCE LISTING ERROR SUMMARY  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 22  
Seq#:4; N Pos. 9,12,18  
Seq#:5; N Pos. 6,12  
Seq#:13; N Pos. 10  
Seq#:13; Xaa Pos. 20,95,99  
Seq#:14; Xaa Pos. 20  
Seq#:17; Xaa Pos. 4  
Seq#:18; N Pos. 3  
Seq#:20; Xaa Pos. 22